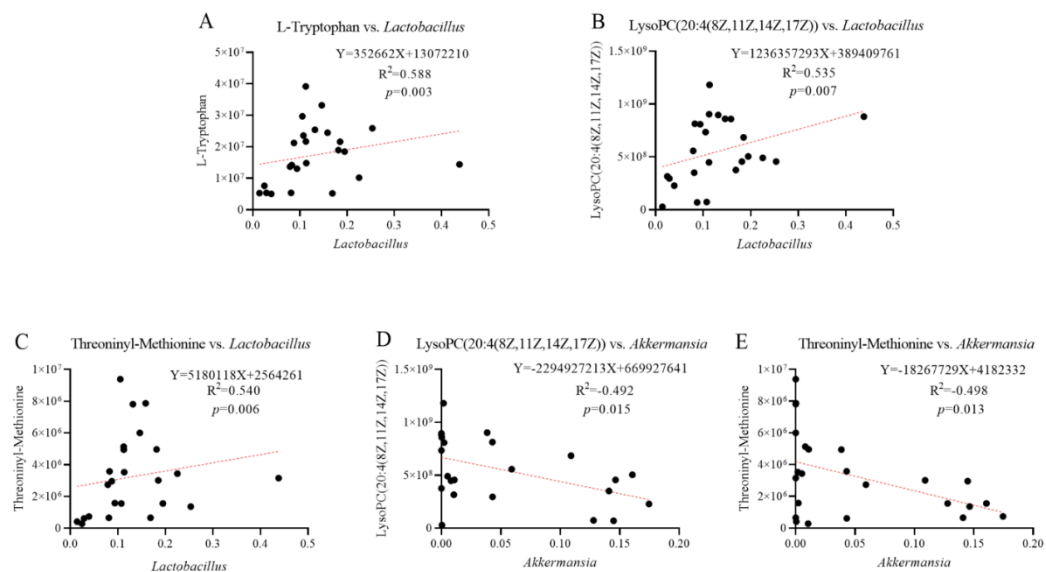
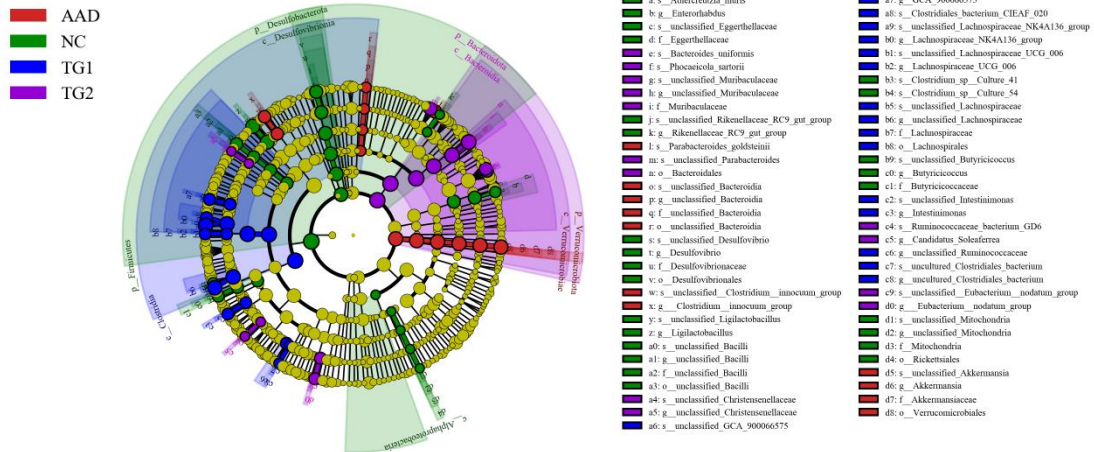


Supplementary Figure S1. Analysis of correlations between cytokine and genus-level intestinal microbes.



Supplementary Figure S2. Analysis of correlations between differential metabolites and genus-level intestinal microbes.

Cladogram



Supplementary Figure S3. LEfSe analysis results of mouse colonic microbiota. The circle radiated from the inside to the outside of the evolutionary branch diagram represents the classification level from gate to species ; each small circle at different classification levels represents a classification at that level. The diameter of the small circle is proportional to the relative abundance. The coloring principle is to uniformly color the species with no significant difference into yellow, and the other different species are colored according to the group with the highest abundance of the species.

Supplementary Table S1. Primer sequences for RT-qPCR detection of colon tissue and amplification of bacterial 16S rRNA genes

Genes/ Region	Forward (5'-3')	Reverse (5'-3')
<i>IL-6</i>	AGCCAGAGTCCTTCAGAGAGA	GCCACTCCTTCTGTGACTCC
<i>IL-1β</i>	AATGCCACCTTTTGACAGTGAT	ATCAGGACAGCCCAGGTCAA
<i>IL-10</i>	CCTGGGTGAGAAGCTGAAGAC	CTTGTAGACACCTTGGTCTTGG
<i>TGF-β</i>	GTGGCTGAACCAAGGAGACG	GTTTGGGGCTGATCCCGTTG
<i>β-actin</i>	GATATCGCTGCGCTGGTCG	CATTCCCACCATCACACCCT
V3-V4	ACTCCTACGGGAGGCAGCA	GGACTACHVGGGTWTCTAAT